

## HISTONE PROPINQUITY USING IMIDOESTERS

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**SUMMARY:** Native chromatin was reacted with bifunctional protein cross-linking reagents of varying molecular sizes. Short cross-linkers do not produce significant yields of polymeric histones whereas intermediate-sized materials such as dimethyladipimide produce extensive interaction between histones. Dimethylsuberimide reacts with all histone fractions but only generates polymers of F<sub>1</sub> and of F<sub>2b</sub> and F<sub>2a2</sub>.

Great interest is currently centered upon chromatin structure and upon the organization of histones upon DNA (1-3). Isolated histone fractions interact together strongly (4,5) and dimethylsuberimide has been utilized to cross-link the components of such histone oligomers (6). We were curious therefore, to see if similar cross-linked products can be obtained from native chromatin, thus providing information about the way in which histones are organized on the DNA molecule. This report describes the use of three bifunctional imidoesters to this end and compares the results with those obtained recently utilizing glutaraldehyde (7, 8).

MATERIALS AND METHODS

Chromatin was isolated as described previously (9). It was adjusted to a concentration of 500  $\mu\text{g/ml}$  DNA and  $5 \times 10^{-4}$  M triethanolamine hydrochloride pH 7.4. Solutions of the imidoesters were prepared immediately before use at 20 mg/ml in the same buffer and the pH adjusted to 7.4 using NaOH. The final concentration of imidoester in the chromatin solution was 1 mg/ml. After fixation for a given time at 4° the reaction mixture was adjusted to 0.4 N H<sub>2</sub>SO<sub>4</sub> and kept in ice for 60 min. The material was centrifuged at 14,000 rpm/90 min in a Sorvall RC-2B centrifuge. The supernatant was dialyzed against ethanol to precipitate histones and polymer histones. Subsequent

analysis was on polyacrylamide gels (10). Gels were scanned in a Beckman Acta III gel scanner.

Suberimide-fixed histones were partially fractionated using the methods of Johns (11).  $F_1$  was extracted in 5% perchloric acid;  $F_{2a}$  and  $F_3$  were extracted in 80% ethanol - 0.25 N HCl, leaving  $F_{2b}$  insoluble.

### RESULTS AND DISCUSSION

The time course for fixation of chromatin by dimethylsuberimide is shown in Fig. 1. The band patterns in acid-urea gels become somewhat blurred as a function of time, no doubt due to the uptake of the cross-linking reagent. There is a rapid decrease in the amount of  $F_1$  histone present, this is followed by a steady decrease in the amount of monomer  $F_{2a2}$  and  $F_{2b}$ . Concurrently we

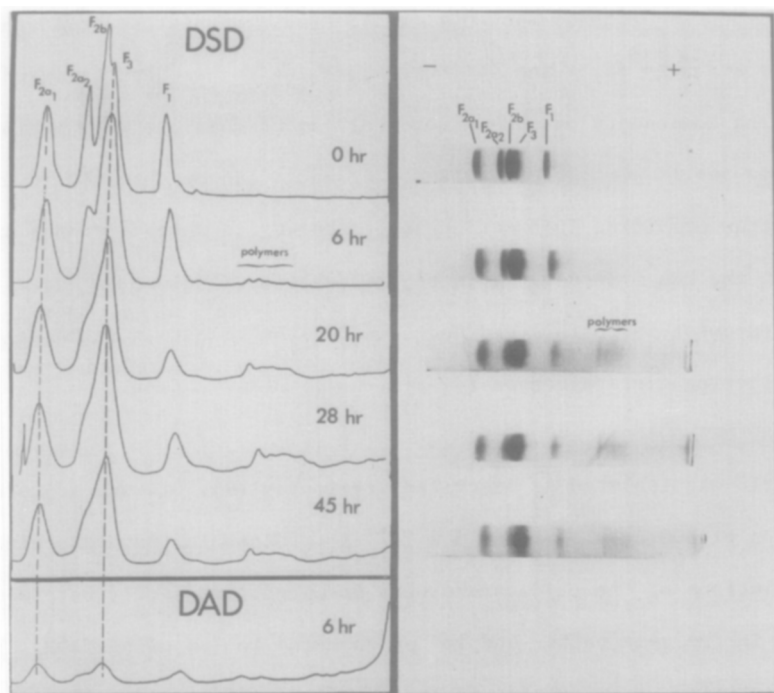


Figure 1. Fixation of histones by dimethylsuberimide on native chromatin. At times indicated the cross-linking reaction was terminated and histones were isolated and analyzed in the acid-urea electrophoretic system (10). The gels (right) show unpolymerized histone as a function of time, together with polymer material migrating primarily in the position normally occupied by dimers. Microdensitometer scans of the gels appear at left with dimethyladipimide fixation after 6 hrs shown for comparison in the bottom panel. The direction of migration is from right to left.

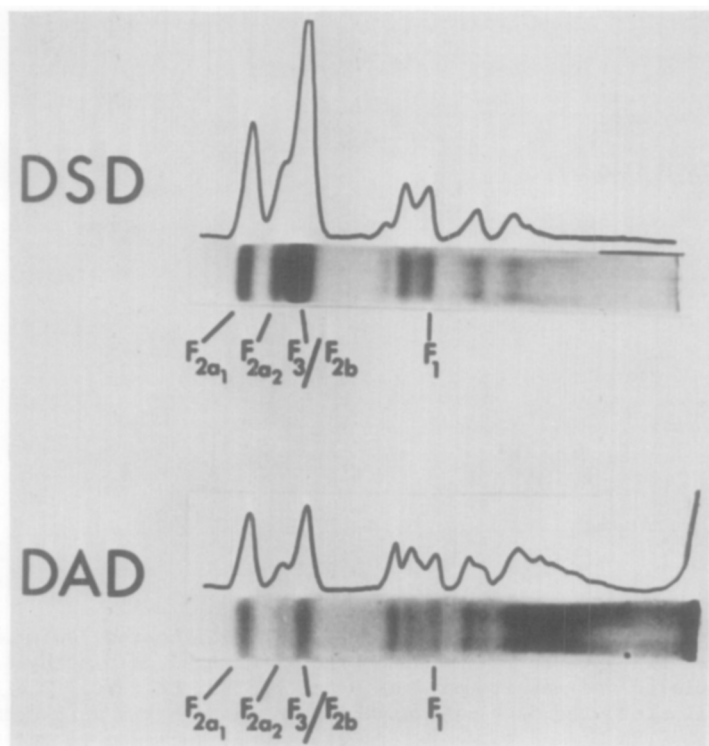


Figure 2. SDS-gel electrophoresis of 28 hr suberimidate-fixed (DSD) and 6 hr adipimidate-fixed (DAD) chromatin. Histone and oligomers were extracted as described in the text, dissolved in SDS-buffer at pH 10.5 and analyzed in the SDS electrophoretic system (10).

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see the appearance of material moving with about 50% of the mobility of monomer histones and at later stages considerably higher molecular weight material is produced. The material with mobility 50% of that of the monomer histone moves in the same region as  $F_3$  dimers and it seems reasonable to ascribe it to various dimers of the histones which are present in ever decreasing amount as the fixation continues. As this material does not build up, it seems likely that it is converted in part to higher polymers as the reaction continues. The gel patterns of Fig. 1 indicate that after 20 hrs a limit is reached and

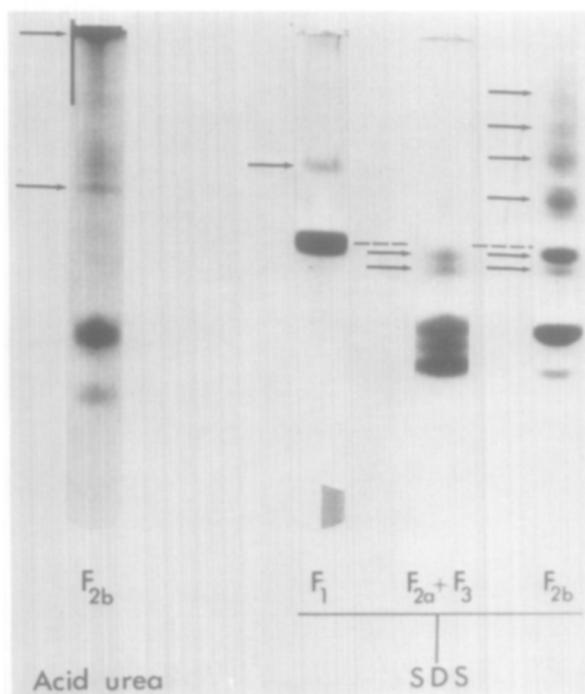


Figure 3. Electrophoretic analysis of partially fractionated, suberimidate-fixed histones. Histones in native chromatin were fixed in dimethylsuberimidate for 20 hrs, isolated and fractionated as described in the text. The  $F_{2b}$  sample (contaminated with 5%  $F_{2a1}$ ) was analyzed in both the acid-urea and SDS gel systems. The  $F_1$  and  $F_{2a} + F_3$  fractions were analyzed in the SDS system. Oligomer bands are denoted with arrows.  $F_1$  in SDS gels migrates more slowly than the dimer bands, as indicated by the dotted line.

this is shown clearly on the microdensitometer traces. During the first 28 hrs of the reaction about 70% of the  $F_1$ ,  $F_{2b}$  and  $F_{2a2}$  was polymerized and is not present in the monomer region, whereas  $F_{2a1}$  and  $F_3$  appear to have undergone relatively little cross-linking reaction. When the reaction was continued for 45 hrs relatively little change was noted, though a small fraction of  $F_{2a1}$  and  $F_3$  has been removed at this time. The products from the polymerization of  $F_{2b}$ ,  $F_1$  and  $F_{2a2}$  may be noted as slower migrating fractions on these scans. This material is highly complex and as many as 6-7 bands may be detected.

Utilizing dimethyladipimidate instead of the suberimidate reveals a remarkable difference. All histone fractions can be cross-linked and disappear

from the gel, though the order of reaction is similar to that seen for the suberimide. A typical residual pattern after 6 hrs of treatment is shown in Fig. 1. In contrast dimethylmalonimide generates no detectable yield of histone polymers and histone monomers do not disappear from the gel even after extended times of reaction.

The products from the suberimide and adipimide fixations were analyzed on SDS gels and the results are shown in Fig. 2. The gel patterns of the suberimide reaction are particularly interesting. Only three classes of polymer are seen, which by analogy with our recent work on glutaraldehyde fixation we would classify as dimers, trimers and tetramers. The relative proportions are indicated in the scans of Fig. 2. Extraction of  $F_1$  reveals that it may be contributing to the slower-moving band, but otherwise the  $F_1$  appears to be lost as much higher molecular weight material which does not enter the gel, and thus we conclude that much of the polymeric material has its origin in  $F_{2b}$  and  $F_{2a2}$  histones. The gel pattern for adipimide fixation is somewhat similar to that for the suberimide-treated material except for two critical differences. These are (1) many higher molecular weight polymers are now observed so that a continuum of material is seen above the level of the intermediate oligomers, and (2) the complexity of the different levels of oligomer is greater than seen following suberimide fixation.

The limited polymer production after incubation with suberimide is not due to hydrolysis of imidoester, as addition of extra material during the course of reaction did not change the time course of polymerization.

I have utilized the chemical fractionation techniques of Johns (11) to further probe the nature of the polymers produced by reaction of chromatin with dimethylsuberimide. The analysis of the products is shown in Fig. 3 and it is apparent that a minor contribution to one of the higher level polymers is coming from  $F_1$ , that the bulk of the polymers is found in the  $F_{2b}$  fraction and that the entire  $F_{2a}$  and  $F_3$  fraction contains very little polymeric material.

Reaction of chromatin at pH 7.4 with dimethyladipimide leads to complete

polymerization of histones so that after several hours they can no longer be extracted into dilute acid. Since material which is acid-soluble during the course of the reaction contains complex histone polymers, we may conclude that histones are cross-linked together and that either the very high molecular weight polymers become insoluble in acid, or that they also become cross-linked to non-histone proteins or DNA.

Since the reaction of chromatin with dimethylsuberimide follows a simple course, this system may prove much more amenable to detailed chemical analysis of histone propinquity than adipimide or glutaraldehyde. Only three of the five histones are polymerized by this agent, about 75% of each of these three fractions ( $F_1$ ,  $F_{2b}$  and  $F_{2a2}$ ) are involved in cross-linking reactions at a stage when essentially none of  $F_3$  or  $F_{2a1}$  have become cross-linked. The products for this reagent are mostly small oligomers of  $F_{2a2}$  and  $F_{2b}$  and higher molecular weight polymers of  $F_1$  which eventually become insoluble in acid. The tendency to form fairly large polymers of  $F_1$  now appears to be well documented (7, 8). If the oligomers of  $F_{2b}$  and  $F_{2a2}$  are analyzed on a logarithmic scale, they appear to be in groups differing by approximately 14,000 daltons, and we suspect that they are dimers, trimers and tetramers. There is apparently relatively little homopolymer of the type  $(F_{2a2})_n$  as the polymers tend to separate into the  $F_{2b}$  fraction upon chemical separation. Thus we incline to the notion that the oligomers are either a mixture of homopolymers of  $F_{2b}$  and heteropolymers of  $F_{2b}$  and  $F_{2a2}$  or consist primarily of the latter possibility. Obviously  $F_{2a1}$  and  $F_3$  are able to react with imidoesters since they are able to form cross-links either to themselves or to other histone fractions using adipimide (or glutaraldehyde) and we may conclude that the lysine residues in these arginine-rich histones involved in adipimide cross-links are situated in close proximity and the suberimide is simply too large a molecule for correct orientation for reaction. [It is intriguing to note that isolated, free  $F_{2a1}$  and  $F_3$  are readily polymerized by dimethylsuberimide to give mixtures of small oligomers (6).] Thus the arginine-rich his-

tones can become involved in cross-links (either to themselves or to other histones) if the cross-linker is longer than 6 Å (malonimide does not form cross-links) or shorter than 13.5 Å (suberimide). Within this range we find cross-linking by adipimide (10.5 Å) and glutaraldehyde (9 Å). Consistent with this proposal are the recent observations that we obtain no fixation with 4,4'-difluoro-3,3'-dinitrodiphenyl sulfone (14.6 Å) or glyoxal (4.5 Å).

Cross-linking is the ultimate result of a reaction sequence with two different rate constants. On the one hand there is the initial attack of the cross-linking reagent to a lysine residue to form a monofunctional imidoester and this is followed by the attack of the imidoester upon a lysine residue in a nearby histone. If this lysine is appropriately situated, one would expect the second reaction to proceed rapidly. Presumably the reason  $F_{2a1}$  and  $F_3$  are not converted into histone polymers at the same rate as  $F_{2a2}$  and  $F_{2b}$  is because the lysine residues are not appropriately organized and the second reaction is much slower than the first and most of the available lysine residues become bound to monofunctional imidoesters so that ultimately few free amino groups are available for cross-linking. However repeating the suberimide reaction at a much lower input concentration of di-imidoester did not change the relative rates of reaction, nor did the relative amounts of the  $F_{2a2}$ ,  $F_{2b}$  oligomers change significantly.

Since it is generally acknowledged that almost all of the DNA is associated with histone, it follows that the distribution of  $F_1$  in sequences and of  $F_{2b}$  and  $F_{2a2}$  in such a manner that the major products are small oligomers imposes certain constraints upon the organization of  $F_3$  and  $F_{2a1}$ , particularly since the total amount of  $F_{2b}$  and  $F_{2a2}$  quite closely approximates that of  $F_3$  and  $F_{2a1}$ . It would seem to be unlikely that all the  $F_3$  and  $F_{2a1}$  are exclusively in the form of tetramers (or higher oligomers) nor can they exist entirely separate from each other (i.e. interspersed with other histones such as  $F_{2b}$ ,  $F_3$ ,  $F_{2a2}$ ). Thus we may anticipate that  $F_3$  and  $F_{2a1}$  are mostly organized in rather small groups of several of 1, 2, 3 or 4 histones.

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